

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/557,351 A
Source: IFW
Date Processed by STIC: 12/12/06

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 12/18/2006

PATENT APPLICATION: US/10/557,351A

TIME: 10:46:00

Input Set : N:\Crf4\12122006\J557351.raw

Output Set: N:\CRF4\12182006\J557351A.raw

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1 <110> APPLICANT: Takada Pharmaceutical Company Limited
2 <120> TITLE OF INVENTION: Antibody and its use
3 <130> FILE REFERENCE: G05-0070
4 <140> CURRENT APPLICATION NUMBER: US/10/557,351A
5 <141> CURRENT FILING DATE: 2005-11-18
6 <150> PRIOR APPLICATION NUMBER: JP2003-151577
7 <151> PRIOR FILING DATE: 2003-05-28
8 <160> NUMBER OF SEQ ID NOS: 20
10 <210> SEQ ID NO: 1
11 <211> LENGTH: 14
12 <212> TYPE: PRT
13 <213> ORGANISM: Artificial Sequence
14 <220> FEATURE:
15 <223> OTHER INFORMATION: immunogen
16 <400> SEQUENCE: 1
17     Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Cys
18             5                      10
20 <210> SEQ ID NO: 2
21 <211> LENGTH: 14
22 <212> TYPE: PRT
23 <213> ORGANISM: Artificial Sequence
24 <220> FEATURE:
25 <223> OTHER INFORMATION: immunogen
26 <400> SEQUENCE: 2
27     Cys His Thr Val Gly Arg Ala Ala Gly Leu Leu Met Gly Leu
28             5                      10
30 <210> SEQ ID NO: 3
31 <211> LENGTH: 16
32 <212> TYPE: PRT
33 <213> ORGANISM: Artificial Sequence
34 <220> FEATURE:
35 <223> OTHER INFORMATION: immunogen
36 <400> SEQUENCE: 3
37     Cys Ala Ser Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu Trp
38             5                      10                      15
40 <210> SEQ ID NO: 4
41 <211> LENGTH: 23
42 <212> TYPE: PRT
43 <213> ORGANISM: Homo sapiens
44 <400> SEQUENCE: 4
45     Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
46     1             5                      10                      15
47     Ala Gly Leu Leu Met Gly Leu

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48                               20
50 <210> SEQ ID NO: 5
51 <211> LENGTH: 30
52 <212> TYPE: PRT
53 <213> ORGANISM: Homo sapiens
54 <400> SEQUENCE: 5
55     Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
56         1                               5               10               15
57     Ala Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu Trp
58         20                               25               30
60 <210> SEQ ID NO: 6
61 <211> LENGTH: 23
62 <212> TYPE: PRT
63 <213> ORGANISM: Rattus norvegicus
64 <400> SEQUENCE: 6
65     Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
66         1                               5               10               15
67     Ser Gly Leu Leu Met Gly Leu
68         20
70 <210> SEQ ID NO: 7
71 <211> LENGTH: 30
72 <212> TYPE: PRT
73 <213> ORGANISM: Rattus norvegicus
74 <400> SEQUENCE: 7
75     Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
76         1                               5               10               15
77     Ser Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu Trp
78         20                               25               30
80 <210> SEQ ID NO: 8
81 <211> LENGTH: 23
82 <212> TYPE: PRT
83 <213> ORGANISM: Mus musculus
84 <400> SEQUENCE: 8
85     Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
86         1                               5               10               15
87     Ser Gly Leu Leu Met Gly Leu
88         20
90 <210> SEQ ID NO: 9
91 <211> LENGTH: 30
92 <212> TYPE: PRT
93 <213> ORGANISM: Mus musculus
94 <400> SEQUENCE: 9
95     Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
96         1                               5               10               15
97     Ser Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Gln Trp
98         20                               25               30
100 <210> SEQ ID NO: 10
101 <211> LENGTH: 23
102 <212> TYPE: PRT

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103 <213> ORGANISM: Sus scrofa
104 <400> SEQUENCE: 10
105      Trp Tyr Lys His Thr Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
106      1              5              10              15
107      Ala Gly Leu Leu Met Gly Leu
108              20
110 <210> SEQ ID NO: 11
111 <211> LENGTH: 30
112 <212> TYPE: PRT
113 <213> ORGANISM: Sus scrofa
114 <400> SEQUENCE: 11
115      Trp Tyr Lys His Thr Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
116      1              5              10              15
117      Ala Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Met Trp
118              20              25              30
120 <210> SEQ ID NO: 12
121 <211> LENGTH: 14
122 <212> TYPE: PRT
123 <213> ORGANISM: Artificial Sequence
124 <220> FEATURE:
125 <223> OTHER INFORMATION: Biotin-labeled peptide
126 <220> FEATURE:
127 <221> NAME/KEY: MOD_RES
128 <222> LOCATION: 14
129 <223> OTHER INFORMATION: Xaa means biotin-labeled Cys modified with Biotin (Long Arm)
Maleimide
130      (Vector Laboratories).
131 <400> SEQUENCE: 12
W--> 132      Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Xaa
133              5              10
135 <210> SEQ ID NO: 13
136 <211> LENGTH: 14
137 <212> TYPE: PRT
138 <213> ORGANISM: Artificial Sequence
139 <220> FEATURE:
140 <223> OTHER INFORMATION: Biotin-labeled peptide
141 <220> FEATURE:
142 <221> NAME/KEY: MOD_RES
143 <222> LOCATION: 1
144 <223> OTHER INFORMATION: Xaa means biotin-labeled Cys modified with Biotin (Long Arm)
Maleimide
145      (Vector Laboratories).
146 <400> SEQUENCE: 13
W--> 147      Xaa His Thr Val Gly Arg Ala Ala Gly Leu Leu Met Gly Leu
148              5              10
150 <210> SEQ ID NO: 14
151 <211> LENGTH: 16
152 <212> TYPE: PRT
153 <213> ORGANISM: Artificial Sequence
154 <220> FEATURE:
155 <223> OTHER INFORMATION: Biotin-labeled peptide

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156 <220> FEATURE:
157 <221> NAME/KEY: MOD_RES
158 <222> LOCATION: 1
159 <223> OTHER INFORMATION: Xaa means biotin-labeled Cys modified with Biotin (Long Arm)
Maleimide
160      (Vector Laboratories).
161 <400> SEQUENCE: 14
W--> 162      Xaa Ala Ser Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu Trp
163              5              10              15
165 <210> SEQ ID NO: 15
166 <211> LENGTH: 328
167 <212> TYPE: PRT
168 <213> ORGANISM: Homo sapiens
169 <400> SEQUENCE: 15
170      Met Asp Asn Ala Ser Phe Ser Glu Pro Trp Pro Ala Asn Ala Ser Gly
171      1              5              10              15
172      Pro Asp Pro Ala Leu Ser Cys Ser Asn Ala Ser Thr Leu Ala Pro Leu
173              20              25              30
174      Pro Ala Pro Leu Ala Val Ala Val Pro Val Val Tyr Ala Val Ile Cys
175              35              40              45
176      Ala Val Gly Leu Ala Gly Asn Ser Ala Val Leu Tyr Val Leu Leu Arg
177      50              55              60
178      Ala Pro Arg Met Lys Thr Val Thr Asn Leu Phe Ile Leu Asn Leu Ala
179      65              70              75              80
180      Ile Ala Asp Glu Leu Phe Thr Leu Val Leu Pro Ile Asn Ile Ala Asp
181              85              90              95
182      Phe Leu Leu Arg Gln Trp Pro Phe Gly Glu Leu Met Cys Lys Leu Ile
183              100              105              110
184      Val Ala Ile Asp Gln Tyr Asn Thr Phe Ser Ser Leu Tyr Phe Leu Thr
185              115              120              125
186      Val Met Ser Ala Asp Arg Tyr Leu Val Val Leu Ala Thr Ala Glu Ser
187      130              135              140
188      Arg Arg Val Ala Gly Arg Thr Tyr Ser Ala Ala Arg Ala Val Ser Leu
189      145              150              155              160
190      Ala Val Trp Gly Ile Val Thr Leu Val Val Leu Pro Phe Ala Val Phe
191              165              170              175
192      Ala Arg Leu Asp Asp Glu Gln Gly Arg Arg Gln Cys Val Leu Val Phe
193              180              185              190
194      Pro Gln Pro Glu Ala Phe Trp Trp Arg Ala Ser Arg Leu Tyr Thr Leu
195              195              200              205
196      Val Leu Gly Phe Ala Ile Pro Val Ser Thr Ile Cys Val Leu Tyr Thr
197      210              215              220
198      Thr Leu Leu Cys Arg Leu His Ala Met Arg Leu Asp Ser His Ala Lys
199      225              230              235              240
200      Ala Leu Glu Arg Ala Lys Lys Arg Val Thr Phe Leu Val Val Ala Ile
201              245              250              255
202      Leu Ala Val Cys Leu Leu Cys Trp Thr Pro Tyr His Leu Ser Thr Val
203              260              265              270
204      Val Ala Leu Thr Thr Asp Leu Pro Gln Thr Pro Leu Val Ile Ala Ile
205              275              280              285

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206      Ser Tyr Phe Ile Thr Ser Leu Ser Tyr Ala Asn Ser Cys Leu Asn Pro
207      290                      295                      300
208      Phe Leu Tyr Ala Phe Leu Asp Ala Ser Phe Arg Arg Asn Leu Arg Gln
209      305                      310                      315                      320
210      Leu Ile Thr Cys Arg Ala Ala Ala
211                      325
213 <210> SEQ ID NO: 16
214 <211> LENGTH: 984
215 <212> TYPE: DNA
216 <213> ORGANISM: Homo sapiens
217 <400> SEQUENCE: 16
218      atggacaacg cctcgttctc ggagccctgg cccgccaacg catcggggccc ggaccgcggcg 60
219      ctgagctgct ccaacgcgct gactctggcg ccgctgccgg cgccgctggc ggtggctgta 120
220      ccagttgtct acgcggtgat ctgcgccgtg ggtctggcgg gcaactccgc cgtgctgtac 180
221      gtggtgctgc gggcgccccg catgaagacc gtcaccaacc tggtcatcct caacctggcc 240
222      atcgccgacg agctcttcac gctgggtgctg cccatcaaca tcgccgactt cctgctgcgg 300
223      cagtggccct tcggggagct catgtgcaag ctcatcgtgg ctatcgacca gtacaacacc 360
224      ttctccagcc tctacttctt caccgtcatg agcgccgacc gctacctggg ggtggtggcc 420
225      actgcggagt cgcgcggggt ggcggggcgc acctacagcg ccgcgcggcg ggtgagcctg 480
226      gccgtgtggg ggatcgctac actcgctcgt ctgcccttcg cagtcttcgc ccggtagac 540
227      gacgagcagg gccggcgcca gtgcgtgcta gtctttccgc agcccgaggc cttctggtgg 600
228      cgcgcgagcc gcctctacac gctcgtgctg ggcttcgcca tccccgtgtc caccatctgt 660
229      gtcctctata ccacctgct gtgccggctg catgccatgc ggctggacag ccacgccaaag 720
230      gccctggagc gcgccaaaga gcgggtgacc ttcttggtgg tggcaatcct ggcggtgtgc 780
231      ctctctgct ggacgcccta ccacctgagc accgtggtgg cgctcaccac cgacctcccg 840
232      cagacgccgc tggtcacgc tatctctac ttcatcacca gcctgagcta cgccaacagc 900
233      tgcctcaacc ccttctctta cgccttctct gacgccagct tccgcaggaa cctccgccag 960
234      ctgataactt gccgcgcggc agcc 984
236 <210> SEQ ID NO: 17
237 <211> LENGTH: 333
238 <212> TYPE: PRT
239 <213> ORGANISM: Homo sapiens
240 <400> SEQUENCE: 17
241      Met Gln Ala Ala Gly His Pro Glu Pro Leu Asp Ser Arg Gly Ser Phe
242      1                      5                      10                      15
243      Ser Leu Pro Thr Met Gly Ala Asn Val Ser Gln Asp Asn Gly Thr Gly
244      20                      25                      30
245      His Asn Ala Thr Phe Ser Glu Pro Leu Pro Phe Leu Tyr Val Leu Leu
246      35                      40                      45
247      Pro Ala Val Tyr Ser Gly Ile Cys Ala Val Gly Leu Thr Gly Asn Thr
248      50                      55                      60
249      Ala Val Ile Leu Val Ile Leu Arg Ala Pro Lys Met Lys Thr Val Thr
250      65                      70                      75                      80
251      Asn Val Phe Ile Leu Asn Leu Ala Val Ala Asp Gly Leu Phe Thr Leu
252      85                      90                      95
253      Val Leu Pro Val Asn Ile Ala Glu His Leu Leu Gln Tyr Trp Pro Phe
254      100                     105                     110
255      Gly Glu Leu Leu Cys Lys Leu Val Leu Ala Val Asp His Tyr Asn Ile
256      115                     120                     125

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/557,351A

DATE: 12/18/2006
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Output Set: N:\CRF4\12182006\J557351A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; Xaa Pos. 14
Seq#:13; Xaa Pos. 1
Seq#:14; Xaa Pos. 1

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:12; Line(s) 129
Seq#:13; Line(s) 144
Seq#:14; Line(s) 159

VERIFICATION SUMMARY

DATE: 12/18/2006

PATENT APPLICATION: US/10/557,351A

TIME: 10:46:01

Input Set : N:\Cr4\12122006\J557351.raw

Output Set: N:\CRF4\12182006\J557351A.raw

L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0

L:147 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0

L:162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0